



SEQ ID NOiG1

Met Thr Glu Lys Asn Gln Thr Val Ser Glu Phe Val Leu Leu Gly Leu Pro Ile Asp Pro Asp
Gln Arg Asp Leu Phe Tyr Ala Leu Phe Leu Ala Met Tyr Val Thr Thr Ile Leu Gly Asn Leu Leu
Ile Ile Val Leu Ile Gln Leu Asp Ser His Leu His Thr Pro Met Tyr Leu Phe Leu Ser Asn Leu Ser
Phe Ser Asp Leu Cys Phe Ser Ser Val Thr Met Pro Lys Leu Leu Gln Asn Met Gln Ser Gln Val
Pro Ser Ile Pro Tyr Ala Gly Cys Leu Thr Gln Met Tyr Phe Phe Leu Phe Phe Gly Asp Leu Glu Ser
Phe Leu Leu Val Ala Met Ala Tyr Asp Arg Tyr Val Ala Ile Cys Phe Pro Leu His Tyr Thr Thr Ile
Met Ser Pro Lys Leu Cys Phe Ser Leu Leu Val Leu Ser Trp Val Leu Thr Met Phe His Ala Val
Leu His Thr Leu Leu Met Ala Arg Leu Cys Phe Cys Ala Asn Thr Ile Pro His Phe Phe Cys Asp
Met Ser Ala Leu Leu Lys Leu Ala Cys Ser Asp Thr Gln Val Asn Glu Leu Val Ile Phe Ile Met Gly
Gly Leu Ile Leu Val Ile Pro Phe Leu Leu Ile Ile Thr Ser Tyr Ala Arg Ile Val Ser Ser Ile Leu Lys
Val Pro Ser Ala Ile Gly Ile Cys Lys Val Phe Ser Thr Cys Gly Ser His Leu Ser Val Val Ser Leu
Phe Tyr Gly Thr Val Ile Gly Leu Tyr Leu Cys Pro Ser Ala Asn Asn Ser Thr Val Lys Glu Thr Ile
Met Ala Met Met Tyr Thr Val Val Thr Pro Met Leu Asn Pro Phe Ile Tyr Ser Leu Arg Asn Lys
Asp Met Lys Gly Ala Leu Arg Arg Val Ile Cys Arg Lys Lys Ile Thr Phe Ser Val

SEQ ID NOiG2

Asp-Pro-Asp-Gln-Arg-Asp-Cys

SEQ ID NOiG3

Leu-Phe-Leu-Ser-Asn-Leu-Ser-Phe-Ser-Asp-Leu-Cys-Ala

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SEQUENCE LISTING

SEQ ID NOiG1

LENGTHiG313

ORGANISMiGCanis familiaris

SEQUENCEiGP30955

DBSOURCEiGswissprot: locus OLFD_CANFA, accession P30955

SEQ ID NOiG2
LENGTHiG7
TYPEiGPRT
ORGANISMiGCanis familiaris
SEQUENCEiGB1
Asp-Pro-Asp-Gln-Arg-Asp-Cys

SEQ ID NOiG3
LENGTHiG13
TYPEiGPRT
ORGANISMiGCanis familiaris
SEQUENCEiGPb2
Leu-Phe-Leu-Ser-Asn-Leu-Ser-Phe-Ser-Asp-Leu-Cys-Ala

a4

cont

